Page 1 of 5 amB



RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/654,462

TIME: 13:46:06

Input Set : A:\PTO.VSK.txt

- 3 -: 110 > APPLICANT: Jessell, Thomas M.
- Briscoe, James
- Rubenstein, John L.R.
- Sander, Maike
- 8 -120 TITLE OF INVENTION: GENETIC DEMONSTRATION OF REQUIREMENT FOR NKX6.1 AND NKX2.2 IN VENTRAL
 - 9 NEURAON GENERATION
 - 11 ·130 FILE REFERENCE: 0575-62166
 - 13 -: 140 > CURRENT APPLICATION NUMBER: 09/654,462
 - 14 141: CURRENT FILING DATE: 2000-09-01
 - 16 <160> NUMBER OF SEQ ID NOS: 4
 - 18 -: 170> SOFTWARE: PatentIn version 3.1
 - 20 -: 210: SEQ ID NO: 1
 - 21 -: 211> LENGTH: 367
 - 22 (212) TYPE: PRT
 - 23 <213> ORGANISM: Human
 - 25 <:400> SEQUENCE: 1
 - 27 Met Leu Ala Val Gly Ala Met Glu Gly Thr Arg Gln Ser Ala Phe Leu

 - 31 Leu Ser Ser Pro Pro Leu Ala Ala Leu His Ser Met Ala Glu Met Lys
 - 20 25
 - 35 Thr Pro Leu Tyr Pro Ala Ala Tyr Pro Pro Leu Pro Ala Gly Pro Pro
 - 40
 - 39 Ser Ser Ser Ser Ser Ser Ser Ser Ser Pro Ser Pro Pro Leu
 - 55
 - 43 Gly Thr His Asn Pro Gly Gly Leu Lys Pro Pro Ala Thr Gly Gly Leu
 - 75
 - 47 Ser Ser Leu Gly Ser Pro Pro Gln Gln Leu Ser Ala Ala Thr Pro His
 - 85 90
 - 51 Gly Ile Asn Asn Ile Leu Ser Arg Pro Ser Met Pro Val Ala Ser Gly
 - 55 Ala Ala Leu Pro Ser Ala Ser Pro Ser Gly Ser Ser Ser Ser Ser Ser
 - 120 115
 - 59 Ser Ser Ala Ser Ala Ser Ser Ala Ser Ala Ala Ala Ala Ala Ala
 - 135
 - 63 Ala Ala Ala Ala Ala Ser Ser Pro Ala Gly Leu Leu Ala Gly Leu

 - 150 155 67 Pro Arg Phe Ser Ser Leu Ser Pro Pro Pro Pro Pro Pro Gly Leu Tyr
 - 165 170
 - 71 Phe Ser Pro Ser Ala Ala Ala Val Ala Ala Val Gly Arg Tyr Pro Lys
 - 180 185
 - 75 Pro Leu Ala Glu Leu Pro Gly Arg Thr Pro Ile Phe Trp Pro Gly Val
 - 195 200
 - 79 Met Gln Ser Pro Pro Trp Arg Asp Ala Arg Leu Ala Cys Thr Pro His

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80 210 215 220	
83 Gln Gly Ser Ile Leu Leu Asp Lys Asp Gly Lys Arg Lys His Thr Arg	
84 225 230 235 240	
87 Pro Thr Phe Ser Gly Gln Gln Ile Phe Ala Leu Glu Lys Thr Phe Glu	
83 245 250 255	
91 Glm. Thr Lys Tyr Leu Ala Gly Pro Glu Arg Ala Arg Leu Ala Tyr Ser	
92 260 265 270	
95 Leu Gly Met Thr Glu Ser Gln Val Lys Val Trp Phe Gln Asn Arg Arg	
96 275 280 285	
99 Thr Lys Trp Arg Lys Lys His Ala Ala Glu Met Ala Thr Ala Lys Lys	
100 290 295 300	
103 Lys Gln Asp Ser Glu Thr Glu Arg Leu Lys Gly Ala Ser Glu Asn Glu	
104 305 310 315	
107 Glu Glu Asp Asp Asp Tyr Asn Lys Pro Leu Asp Pro Asn Ser Asp Asp	
108 325 330 335	
111 Glu Lys Ile Thr Gln Leu Leu Lys Lys His Lys Ser Ser Ser Gly Gly	
112 340 345 350	
115 Gly Gly Gly Leu Leu His Ala Ser Glu Pro Glu Ser Ser	
116 355 360 365	
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122 -: 1213' ORGANISM: Human	
124 -: (400) SEQUENCE: 2	<i>C</i> 0
125 cgtgggatgt tagcggtggg ggcaatggag ggcacccggc agagcgcatt cctgctcagc	60
127 agoodtooc tggccgcct gcacagcatg gccgagatga agacccgct gtacctgcc	120
129 gcgtatecce cgctgcctge cggccccce tectectegt cetegtegte gteetecteg	180 240
131 tegecetece egectetggg cacceacaac ceaggeggee tgaageecee ggecaegggg	300
133 gggeteteat eceteggeag eeceeggag eagetetegg eegeeacee acaeggeate	360
135 aacaatatee tgageeggee etecatgeee gtggeetegg gggeegeeet geeeteegee	420
137 tegeceteeg gitecteete eteetettee tegicegeet etgecteete egectetgee	480
139 geogeogogg etgetgeege ggeogoagee geogeoteat eeceggeggg getgetggee 141 ggaetgeeae getttageag eetgageeeg eegeegeege egeooggget etaetteage	540
	600
143 cocagogog eggeegtgge egeegtggge eggtaceeea ageegetgge tgagetgeet 145 ggeeggaege eeatettetg geeeggagtg atgeagagee egeeetggag ggaegeaege	660
143 gyddyddyd ddaididdy gdddyddig alyddydydd dydddigyd gyddyddigiol 147 ciggodigia ocociogiga gi	682
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151 (211) LENGTH: 185	
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158 ctttttccgg acagcagate ttegecetgg agaagacttt egaacaaaca aaatacttgg	120
160 eggggeega gagggetegt tiggeetatt egitiggggat gacagagagt caggicaagg	180
162 tqagt	185
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167 :1212 TYPE: DNA	
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175	gaggaagagg	acgacgacta	caataagcct	ctggatccsa	actcggacga	cgagaaaatc	180
177	acgcagetgt	tgaagaagca	caagtccagc	agcggcggcg	geggeggeet	cctactgcac	240
179	gcgtccgagc	cggagagctc	atcctgaacg	ccg			273

VERIFICATION SUMMARY

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